

Result	8	Query	Match	Length	DB	ID	Description
No.	Score						
c	1	508.6	37.3	676	10	BE385990	BB385990 601276636
c	2	503.6	37.0	591	10	AT1884686	AT1884686 w183d07.x
c	3	502.4	36.9	534	10	BE350014	BB350014 ht07912.x
c	4	494.8	36.3	814	10	AL567376	AL567376
c	5	495.8	35.2	578	10	AT1742092	AT1742092 wq38h03.x
c	6	471.2	34.6	759	11	BR126050	BR126050 601176288
c	7	464	34.1	843	11	BF125134	BF125134 601762356
c	8	429.6	31.5	478	10	AT136826	AT136826 wp69h10.x
c	9	413.6	30.4	460	10	AI1937602	AI1937602 wp81f07.x
c	10	409.9	30.0	690	10	AW149665	AW149665 xf40h05.x
c	11	406.6	29.9	821	11	BG152229	BG152229 602731372
c	12	386	28.3	387	10	AI272281	AI272281 ap23c03.x

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

ALIGNMENTS

REFERENCES

AUTHORS

VERSION

KEYWORDS

SOURCE

ORGANISM

COMMENT

Email: cgabs-r@mail.nih.gov

Tissue Procurement: ARCC/DCRD/DP

CDNA Library Preparation: Ling Hong/Rubin Laboratory

CDNA Library Arrayed by: The L.M.A.G.E. Consortium (LLNL)

DNA Sequencing by: Incyte Genomics, Inc.

Clone distribution: MGC clone distribution information can be found through the L.M.A.G.E. Consortium/LLNL at: image.llnl.gov

Plate: ILCM285 Row: y Column: 09

High quality sequence stop: 653.

FEATURES

Source

1. .676

/organism="Homo sapiens"

/db_xref="taxon:9606"

/clone "IMAGE_3617696"

/clone_1lb="NIH_MGC_20"

/issue_type="melanotic melanoma"

/lab_host="DH10B (phage-resistant)"

/note="Organ: skin; Vector: pOTB7; Site_1: XbaI; Site_2:

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DEFINITION		IMAGE:2367413					
REFERENCE		1 (bases 1 to 578)	similar to	SP-6P39_HUMAN	043194	PUTATIVE G	
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1. . 759

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ECORI; cDNA made by oligo-dT priming. Directionally

cloned into EcoRI/XbaI sites using the following 5'

adaptor: GGCACGAG(G). Size-selected >500bp for average

insert size 1.8kb. Library constructed by Ling Hong in

the laboratory of Gerald M. Rubin (University of

California, Berkeley) using ZAP-cDNA synthesis kit

(Stratagene) and superscript II RT (Life Technologies)."

BASE COUNT

191 a

192 c

221 g

155 t

ORIGIN

AUTHORS NIH-MGC http://mgc.nci.nih.gov/.

TITLE National Institutes of Health, Mammalian Gene Collection (MGC)

JOURNAL Unpublished (1999)

COMMENT

Email: cgbabs-r@mail.nih.gov

Tissue Procurement: ATCC/ATCC/ATCC

CDNA Library Preparation: Ling Hong/Rubin Laboratory

CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LNL)

DNA Sequencing by: Incyte Genomics, Inc.

Clone distribution: MGC clone distribution information can be

found through the I.M.A.G.E. Consortium/LNL at: Image.lnl.gov

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High quality sequence stop: 711

Location/Qualifiers

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/note="Organ: skin; Vector: pMB7; Site_1: XbaI; Site_2:

ECORI; cDNA made by oligo-dT priming. Directionally

cloned into EcoRI/XbaI sites using the following 5'

adaptor: GGCACGAG(G). Size-selected >500bp for average

insert size 1.8kb. Library constructed by Ling Hong in

the laboratory of Gerald M. Rubin (University of

California, Berkeley) using ZAP-cDNA synthesis kit

(Stratagene) and superscript II RT (Life Technologies)."

BASE COUNT

225 a

192 c

255 g

171 t

ORIGIN

RESULT

7

BF125134/c

LOCUS

BP125134

DEFINITION

601762356F1 NIH_MGC_20 Homo sapiens cDNA clone IMAGE:4025340

mRNA sequence.

ACCESSION

BR125134

VERSION

BR125134.1

SOURCE

EPR.

ORGANISM

Homo sapiens

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

REFERENCE

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Wed Jan 9 08:51:39 2002

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